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ID 091A69 PRELIMINARY: PRT; 197 AA.
AC 091A69;
DT 01-OCT-2000 (Tremblrel. 15, Created)
DT 01-OCT-2000 (Tremblrel. 15, Last sequence update)
DT 01-JUN-2001 (Tremblrel. 17, Last annotation update)
DE INHIBITOR OF APOPTOSIS 1 (FRAGMENT).
OS Gallus gallus (Chicken).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Archosauria; Aves; Neognathae; Galliformes; Phasianidae; Phasianinae;
OC Gallus.
NCBI_TaxID=9031;
RN [1]
RP SEQUENCE FROM N.A.
RC STRAIN-BRED FAYOUMI; TISSUE=SPLEEN;
RA Zhou H, Lamont S.J.;
RT *Genetic variation among chicken lines and mammalian species in
RT specific genes.
RL Submitted (JAN-2000) to the EMBL/GenBank/DBJ databases.
DR EMBL: AF221083; AAF35320.1; -.
DR InterPro: IPR001370; BIR.
DR Pfam: PF00653; BIR; 2.
DR SMART: SM00238; BIR; 2.
DR PROSITE: PS01282; BIR_REPEAT_1; 2.
DR PROSITE: PS0143; BIR_REPEAT_2; 2.
FT NON_TER 1 1
FT 197 197
SQ SEQUENCE 197 AA; 22602 MW; D7923DAACFE23E1A CRC64;

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Query Match 98.3%; Score 290; DB 13; Length 197;
Best Local Similarity 95.8%; Pred. No. 6,3e-30;
Matches 46; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

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QY 1 PQLASAGFYVGRNDVVCFCDCGLRCWESGDDPVVHAHAKFPFCE 48
DB 135 PQLADAGFYVGRNDVVCFCDCGLRCWESGDDPVVHAHAKFPFCE 182

RESULT 3
057319 PRELIMINARY: PRT; 610 AA.
AC 057319;
DT 01-JUN-1998 (Tremblrel. 06, Created)
DT 01-JUN-1998 (Tremblrel. 06, Last sequence update)
DT 01-JUN-2001 (Tremblrel. 17, Last annotation update)
DE INHIBITOR OF APOPTOSIS PROTEIN 1 (IAP) (INHIBITOR OF T CELL APOPTOSIS
DE PROTEIN).
GN IAP1.
OS Gallus gallus (Chicken).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Archosauria; Aves; Neognathae; Galliformes; Phasianidae; Phasianinae;
OC Gallus.
NCBI_TaxID=9031;
RN [1]
RP SEQUENCE FROM N.A.
RC STRAIN-WHITE LEGHORN; TISSUE=EMBRYONIC FIBROBLAST;
RX MEDLINE=98038601; Pubmed=9372964;
RA You M, Ku P.-T., Hrdlickova R., Bose H.R. Jr.;
RT *Ch-IAP1, a member of the inhibitor-of-apoptosis protein family, is a
RT mediator of the antiapoptotic activity of the v-rel oncoprotein.
RL Mol. Cell. Biol. 17:7328-7341(1997).
CC -1- FUNCTION: SUPPRESSOR OF APOPTOSIS IN ONCOPROTEIN V-REL-TRANSFORMED
CC CELLS.
CC -1- SUBCELLULAR LOCATION: EXPRESSED PREDOMINANTLY IN THE CYTOPLASM OF
CC THE V-REL-TRANSFORMED CELLS.
CC -1- TISSUE SPECIFICITY: EXPRESSED AT RELATIVELY HIGH LEVELS IN THE
CC SPLEEN, THYMUS, BURSA, INTESTINE, AND LUNG, AND AT VERY LOW LEVELS
CC IN TESTIS, BRAIN, AND SKELETAL MUSCLE.
CC -1- INDUCTION: INDUCED DURING THE V-REL-MEDIATED TRANSFORMATION
CC PROCESS.
CC -1- DOMAIN: THE RING FINGER IS IMPORTANT FOR ITS ANTIAPOPTOTIC EFFECT.
CC -1- SIMILARITY: MEMBER OF THE IAP FAMILY.
CC -1- SIMILARITY: CONTAINS 3 BIR DOMAINS (BACULOVIRAL INHIBITION OF

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CC APOPTOSIS PROTEIN REPEAT).
CC -1- SIMILARITY: CONTAINS A C3HC4-CLASS ZINC FINGER.
DR EMBL: AF008592; AAB88044.1; -.
DR HSSP: Q13490; IOBH.
DR InterPro: IPR001370; BIR.
DR InterPro: IPR001315; CARD.
DR InterPro: IPR001841; Znf_ring.
DR Pfam: PF00653; BIR; 3.
DR Pfam: PF00619; CARD; 1.
DR SMART: SM00097; zf-C3HC4; 1.
DR SMART: SM00238; BIR; 3.
DR SMART: SM00114; CARD; 1.
DR SMART: SM00184; RING; 1.
DR PROSITE: PS01282; BIR_REPEAT_1; 3.
DR PROSITE: PS0143; BIR_REPEAT_2; 3.
DR PROSITE: PS0209; CARD; 1.
KW Apoptosis; Zinc-finger; Repeat.
FT REPEAT 30 97 BIR REPEAT 1.
FT REPEAT 176 242 BIR REPEAT 2.
FT REPEAT 262 329 BIR REPEAT 3.
FT ZN_FING 563 597 C3HC4-TYPE.
SQ SEQUENCE 610 AA; 68924 MW; ADF47619650B44A6 CRC64;

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Query Match 98.3%; Score 290; DB 13; Length 610;
Best Local Similarity 95.8%; Pred. No. 2e-29;
Matches 46; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

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QY 1 PQLASAGFYVGRNDVVCFCDCGLRCWESGDDPVVHAHAKFPFCE 48
DB 280 PQLADAGFYVGRNDVVCFCDCGLRCWESGDDPVVHAHAKFPFCE 327

RESULT 4
09UNH1 PRELIMINARY: PRT; 1140 AA.
AC 09UNH1;
DT 01-MAY-2000 (Tremblrel. 13, Created)
DT 01-MAY-2000 (Tremblrel. 13, Last sequence update)
DT 01-JUN-2001 (Tremblrel. 17, Last annotation update)
DE APl2-MLT FUSION PROTEIN.
GN APl2-MLT.
OS Homo sapiens (Human).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Mammalia; Eutheria; Primates; Catarrhini; Homiidae; Homo.
OX NCBI_TaxID=9606;
RN [1]
RP SEQUENCE FROM N.A.
RX MEDLINE=99272400; Pubmed=10339464;
RA Dierlamm J., Baens M., Wlodarska I., Stefanova-Uzunova M.,
RA Hernandez J.M., Hossfeld D.K., De Wolf-Peters C., Hagemeijer A.,
RA Van den Bergh H., Marynen P.;
RT *The apoptosis inhibitor gene APl2 and a novel 189 gene, MLT, are
RT recurrently rearranged in the t(11;18)(q21;q21)p65associated with
RT BLOOD 93:3601-3609(1999).
CC -1- SIMILARITY: TO IMMUNOGLOBULIN AND MAJOR HISTOCOMPATIBILITY COMPLEX
CC -1- DOMAIN.
DR EMBL: AF123094; AAD46161.1; -.
DR HSSP: Q13490; IOBH.
DR InterPro: IPR001370; BIR.
DR InterPro: IPR003576; Caspase.
DR InterPro: IPR001309; ICE_p20.
DR InterPro: IPR003598; IC_c2.
DR InterPro: IPR003600; IC_c1.
DR InterPro: IPR003006; IC_MHC.
DR Pfam: PF00653; BIR; 3.
DR Pfam: PF00047; IC; 2.
DR SMART: SM00238; BIR; 3.
DR SMART: SM00115; CARD; 1.
DR SMART: SM00408; IC_c2; 1.
DR SMART: SM00410; IC_c1; 1.
DR PROSITE: PS01282; BIR_REPEAT_1; 3.

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DR PROSITE: PS50143; BIR_REPEAT_2; 3.
DR PROSITE: PS50208; CASPASE_P20; 1.
SQ SEQUENCE 1140 AA; 128738 MW; 0C18D890287C723E CRC64;

Query Match 95.6%; Score 282; DB 4; Length 1140;
Best Local Similarity 93.8%; Pred. No. 4, 2e-28;
Matches 45; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

OY 1 PEOIASAGFYVGRNDYKCFCCDGLRCWESGDDPWVHAHAKFPCE 48
|||||
DB 273 PEOIASAGFYVGRNDYKCFCCDGLRCWESGDDPWVHAHAKFPCE 320

RESULT 5
O9QZC6 PRELIMINARY; PRT; 589 AA.

AC O9QZC6 (TREMBLrel. 13, Created)
DT 01-MAY-2000 (TREMBLrel. 13, last sequence update)
DE 01-JUN-2001 (TREMBLrel. 17, last annotation update)
DE INHIBITOR OF APOPTOSIS PROTEIN 2.
OS Rattus norvegicus (Rat).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Rattus.
OX NCBI_TaxID=10116;
RN [1]
RP SEQUENCE FROM N.A.
RC TISSUE=KIDNEY;
RA Dong Z., Denton M., Gu S.M., Salkumar P., Venkathalam M.A.;
RT "Cloning of cDNA for rat inhibitor of apoptosis protein 2.";
RL Submitted (SEP-1999) to the EMBL/GenBank/DBJ databases.
CC -1- SIMILARITY: CONTAINS A RING-TYPE ZINC FINGER.
DR EMBL: AF190020; AAF04585.1; -
DR HSSP: Q13490; 108H.
DR InterPro: IPR001370; BIR.
DR InterPro: IPR001315; CARD.
DR Pfam: PF00653; BIR; 3
DR Pfam: PF00619; CARD; 1.
DR Pfam: PF00097; ZF-C3HC4; 1.
DR SMART: SM00238; BIR; 3.
DR SMART: SM00114; CARD; 1.
DR SMART: SM00184; RING; 1.
DR PROSITE: PS01282; BIR_REPEAT_1; 3.
DR PROSITE: PS50143; BIR_REPEAT_2; 3.
DR PROSITE: PS50209; CARD; 1.
KM ZINC FINGER.
SQ SEQUENCE 589 AA; 66777 MW; E6812FE3EA34142 CRC64;

Query Match 93.9%; Score 277; DB 11; Length 589;
Best Local Similarity 91.7%; Pred. No. 9, 4e-28;
Matches 44; Conservative 1; Mismatches 3; Indels 0; Gaps 0;

OY 1 PEOIASAGFYVGRNDYKCFCCDGLRCWESGDDPWVHAHAKFPCE 48
|||||
DB 259 PEOIASAGFYVGRNDYKCFCCDGLRCWESGDDPWVHAHAKFPCE 306

RESULT 6
O9ESE8 PRELIMINARY; PRT; 589 AA.

AC O9ESE8 (TREMBLrel. 16, Created)
DT 01-MAR-2001 (TREMBLrel. 16, last sequence update)
DT 01-JUN-2001 (TREMBLrel. 17, last annotation update)
DE INHIBITOR OF APOPTOSIS PROTEIN 2.
OS Rattus norvegicus (Rat).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Rattus.
OX NCBI_TaxID=10116;
RN [1]

RP SEQUENCE FROM N.A.
RA Holcik M., Lefebvre C.A., Hicks K., Korneluk R.G.;
RT "Cloning and Characterization of the Rat Homologs of the Inhibitor of
RT Apoptosis Protein 1, 2, and 3 Genes.";
RL Submitted (SEP-1999) to the EMBL/GenBank/DBJ databases.
CC -1- SIMILARITY: CONTAINS A RING-TYPE ZINC FINGER.
DR EMBL: AF183431; AAC22971.1; -
DR InterPro: IPR001370; BIR.
DR InterPro: IPR001315; CARD.
DR InterPro: IPR001841; ZNF_Fing.
DR Pfam: PF00653; BIR; 3.
DR Pfam: PF00619; CARD; 1.
DR Pfam: PF00097; ZF-C3HC4; 1.
DR SMART: SM00238; BIR; 3.
DR SMART: SM00114; CARD; 1.
DR SMART: SM00184; RING; 1.
DR PROSITE: PS01282; BIR_REPEAT_1; 1.
DR PROSITE: PS50143; BIR_REPEAT_2; 3.
DR PROSITE: PS50209; CARD; 1.
KM ZINC FINGER.
SQ SEQUENCE 589 AA; 66750 MW; B4F7089BD7CD285B CRC64;

Query Match 93.9%; Score 277; DB 11; Length 589;
Best Local Similarity 91.7%; Pred. No. 9, 4e-28;
Matches 44; Conservative 1; Mismatches 3; Indels 0; Gaps 0;

OY 1 PEOIASAGFYVGRNDYKCFCCDGLRCWESGDDPWVHAHAKFPCE 48
|||||
DB 259 PEOIASAGFYVGRNDYKCFCCDGLRCWESGDDPWVHAHAKFPCE 306

RESULT 7
O9ESE9 PRELIMINARY; PRT; 602 AA.

AC O9ESE9 (TREMBLrel. 16, Created)
DT 01-MAR-2001 (TREMBLrel. 16, last sequence update)
DT 01-JUN-2001 (TREMBLrel. 17, last annotation update)
DE 01-JUN-2001 (TREMBLrel. 17, last annotation update)
DE INHIBITOR OF APOPTOSIS PROTEIN 1.
OS Rattus norvegicus (Rat).
OC Rattus norvegicus (Rat).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Rattus.
OX NCBI_TaxID=10116;
RN [1]
RP SEQUENCE FROM N.A.
RA Holcik M., Lefebvre C.A., Hicks K., Korneluk R.G.;
RT "Cloning and Characterization of the Rat Homologs of the Inhibitor of
RT Apoptosis Protein 1, 2, and 3 Genes.";
RL Submitted (SEP-1999) to the EMBL/GenBank/DBJ databases.
CC -1- SIMILARITY: CONTAINS A RING-TYPE ZINC FINGER.
DR EMBL: AF183430; AAC22970.1; -
DR InterPro: IPR001370; BIR.
DR InterPro: IPR001315; CARD.
DR InterPro: IPR001841; ZNF_Fing.
DR Pfam: PF00653; BIR; 3.
DR Pfam: PF00619; CARD; 1.
DR Pfam: PF00097; ZF-C3HC4; 1.
DR SMART: SM00238; BIR; 3.
DR SMART: SM00114; CARD; 1.
DR SMART: SM00184; RING; 1.
DR PROSITE: PS01282; BIR_REPEAT_1; 1.
DR PROSITE: PS50143; BIR_REPEAT_2; 3.
DR PROSITE: PS50209; CARD; 1.
KM ZINC FINGER.
SQ SEQUENCE 602 AA; 67326 MW; CC91385FEA62DE5A CRC64;

Query Match 93.2%; Score 275; DB 11; Length 602;
Best Local Similarity 89.6%; Pred. No. 1, 7e-27;
Matches 43; Conservative 3; Mismatches 2; Indels 0; Gaps 0;

OY 1 PEOIASAGFYVGRNDYKCFCCDGLRCWESGDDPWVHAHAKFPCE 48

Db 273 PDELASAGFYTTGHSDDVKFCDDGGLRCWESGDDPWEHAKWFPCE 320

RESULT	6
09DDN2	
ID	09DDN2
AC	09DDN2; PRELIMINARY; PRT: 324 AA.
DT	01-MAR-2001 (TREMBLrel. 16, Created)
DT	01-MAR-2001 (TREMBLrel. 16, last sequence update)
DT	01-JUN-2001 (TREMBLrel. 17, last annotation update)
DE	APOPTOSIS INHIBITOR CH-1A1 (FRAGMENT).
OS	Gallus gallus (Chicken).
OC	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC	Archosauirae; Aves; Neognathae; Galliformes; Phasianidae; Phasianinae;
OC	Gallus.
OX	NCBI_TaxId=9031;
RN	[1]
RP	SEQUENCE FROM N.A.
RA	Pendleton C.N., Bergmann W.J., Varadarajan J., Bose H.R., Jr.:
RT	"The apoptosis inhibitor ch-1a1 is a direct transcriptional target of
RT	v-Rel and c-Rel."
RL	Submitted (OCF-2000) to the EMBL/Genbank/DBJ databases.
DR	EMBL: AF311289; AAC42316.1; -
DR	InterPro: IPR001370; BIR.
DR	Pfam: PF00653; BIR: 3
DR	SMART: SM00238; BIR: 3
DR	PROSITE: PS01282; BIR_REPEAT_1; 1
DR	PROSITE: PS01433; BIR_REPEAT_2; 3.
DR	NON_TER 324 324
EQ	SEQUENCE 324 AA; 36567 MW; 5E2B89DEAE373F35 CRC64;

Query Match	91.9%;	Score 271;	DB 13;	Length 324;
Best Local Similarity	95.68;	Pred. No. 3e-27;		
Matches 43; Conservative	1;	Mismatches 1;	Indels 0;	Gaps 0;

Qy 1 PEQLASAGFYVVGRRNDVKCFCCDGLRCWESGDDPWVEHAKWFP 45
 ||||| ||||||| : |||||
Dd 280 PEQLADAGFYVVGRRNDVKCFCCDGLRCWESGDDPWIEHAKWFP 324

RESULT	9	
088642		
ID	088642:	PRELIMINARY: PRT; 224 AA.
DT	01-NOV-1998 (TREMBLrel. 08.	Created)
DT	01-NOV-1998 (TREMBLrel. 08.	Last sequence update)
DT	01-JUN-2001 (TREMBLrel. 17.	Last annotation update)
DE	INHIBITOR OF APOPTOSIS PROTEIN (FRAGMENT).	
GN	RIAP1.	
OS	Rattus norvegicus (Rat); Chordata; Craniata; Vertebrata; Euteleostomi;	
OC	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;	
OC	Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Rattus	
OX	NCBI_TaxID=10116;	
RN	RN [1]	
RP	SEQUENCE FROM N.A.	
RC	STRAIN=MISTAR; TISSUE=OVARY, CORPUS LUTEUM;	
RA	Bradley C.R., Latreu R.R., Dharmarajan A.M.;	
RT	Cloning and characterisation of an inhibitor of apoptosis protein	
RT	(IAP) In the rat corpus luteum."	
RL	Submitted (Aug-1998) to the EMBL/GenBank/DDA databases.	
DR	EMBL: AF081503; AAC32497.1;	
DR	HSSP: 013490; 108H.	
DR	InterPro: IP0001370;	BIR.
DR	Pfam: PF00653;	BIR; 2.
DR	SMART: SM00238;	BIR; 2.
DR	PROSITE: PS50143;	BIR_REPEAT_2; 3.
FT	NON_TER	1
FT	NON_TER	224
FT	NON_TER	224
QO	SEQUENCE	224 AA; 25209 MW; 213A5253AD5EB56A CRC64;

Query Match	70.88;	Score 209;	DB 11;	Length 224;
Best Local Similarity	91.98;	Pred. No. 2.2e-19;		
Matches 34; Conservative	0;	Mismatches 3;	Indels 0;	Gaps 0;

QY 1 PEOLASAGFYVYVGRNDVDKFCDCDGLRCWESGDDPW 37
 |||||
 Db 188 PEOLASAGFYVDHNDVDKFCDCDGLRCWEGGDDPW 222

RESULT	10
09EN27	
ID 09EN27	PRELIMINARY; PRT; 264 AA.
AC 09EN27	
DT 01-MAR-2001 (TREMBLrel. 16, Created)	
DT 01-MAR-2001 (TREMBLrel. 16, Last sequence update)	
DT 01-JUN-2001 (TREMBLrel. 17, Last annotation update)	
DE AMV021.	
GN AMV021.	
OS Amsacta moorei entomopoxvirus (AmpPV).	
OC Viruses; dsDNA viruses, no RNA stage; Poxviridae; Entomopoxvirinae	
OC Entomopoxvirus B.	
OX NCBI_TaxId=28321;	
RN [1]	
RP SEQUENCE FROM N.A.	
RX MEDLINE=20396580; PubMed=10936094;	
RA Bawden A.L., Glassberg K.J., Diggins J., Shaw R., Farmerie W.,	
RA Moyer R.W.;	
RT "Complete Genomic Sequence of the Amsacta moorei Entomopoxvirus:	
RT Analysis and Comparison with Other Poxviruses.";	
RL Virology 274:120-139(2000).	
RN [2]	
RP SEQUENCE FROM N.A.	
RA Bawden A.L., Glassberg K.J., Diggins J., Shaw R., Farmerie W.,	
RA Moyer R.W.;	
RL Submitted (MAR-2000) to the EMBL/GenBank/DBJ databases.	
EMBL AF250284; AACG2727.1; -	
DR InterPro: IPR001370; BIR.	
DR InterPro: IPR001841; znf_r10g.	
DR Pfam: PR00653; BIR. 2	
DR SMART: SM00238; BIR. 2	
DR SMART: SM00184; RING. 1.	
DR PROSITE: PS50143; BIR.REPEAT_2; 2.	
DR SSQSDNCE 264 AA; 30547 MW; 2EB2DA4B58D920A CRC64;	

Query Match	67.1%;	Score 198;	DB 12;	Length 264;
Best Local Similarity	61.7%;	Pred. No. 7e-18;		
Matches 29;	Conservative 10;	Mismatches 8;	Indels 0;	Gaps 0;

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QY      2  EQLASAGFYVVGRRDDVKCFCCDGGRLRCWESGDDPWVEHAKWFECE 48
      1 : : : 1 : : : 1 : : : 1 : : : 1 : : : 1 : : : 1 : : :
Db      126 EKLAEGAFYTGKSDKVKCFYCDGGLNKWETDDDDPWIQHARWFDKCD 172
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RESULT	11	
09HAP7		
ID	09HAP7	PRELIMINARY; PRT; 280 AA.
AC	09HAP7	
DT	01-MAR-2001 (TREMBL)	16, Created)
DT	01-MAR-2001 (TREMBL)	16, Last sequence update)
DT	01-JUN-2001 (TREMBL)	17, Last annotation update)
DE	LIVIN INHIBITOR OF APOPTOSIS (INHIBITOR OF APOPTOSIS) (BA261N1.1.1.1)	
DE	(BACULOVIRAL IAP REPEAT-CONTAINING PROTEIN 7 (LIVIN), ISOFORM 1).	
OS	LIVIN OR BIRC7.	
GN	Homo sapiens (Human).	
OC	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;	
OC	Mammalia; Eulipotyria; Primates; Carnivora; Hominoidea; Homo.	
OX	NCBI_Taxid=9606;	
OX	[1]	
RN	SEQUENCE FROM N.A.	
RP	Kasof G.M., Gomes B.C.;	
RA	"Livin, a novel inhibitor of apoptosis (IAP) family member.";	
RT	J. Biol. Chem. 0:0-0(2000).	

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RN [12]
RP SEQUENCE FROM N.A.
RA Ashab Y., Allan A., Pollack A., Panet A., Ben-Yehuda D.;
RT "Two splicing variants of a new inhibitor of apoptosis gene with
RL different biological properties and tissue distribution pattern.";
RN Submitted (Apr-2001) to the EMBL/GenBank/DBJ databases.
[13]
RP SEQUENCE FROM N.A.
RA Hall R.;
RL Submitted (Apr-2001) to the EMBL/GenBank/DBJ databases.
CC -1- SIMILARITY: CONTAINS A RING-TYPE ZINC FINGER.
DR EMBL; AF311388; AAC33622.1; -
DR EMBL; AJ309298; CAC37337.1; -
DR EMBL; AL121827; CAC36112.1; -
DR InterPro: IPR001370; BIR.
DR InterPro: IPR001841; Znf_fing.
DR Pfam: PF00653; BIR; 1.
DR Pfam: PF00097; zf-C3HC4; 1.
DR SMART; SM00238; BIR; 1.
DR SMART; SM00184; RING; 1.
DR PROSITE; PS50143; BIR_REPEAT_2; 1.
DR PROSITE; PS00518; ZINC_FINGER_C3HC4; UNKNOWN_1.
DR Zinc-finger.
SQ SEQUENCE 280 AA; 30866 MW; 630BE9C0737F7952 CRC64;

Query Match 66.1%; Score 195; DB 4; Length 280;
Best Local Similarity 62.5%; Pred. No. 1.9e-17;
Matches 30; Conservative 7; Mismatches 11; Indels 0; Gaps 0;

QY 1 PEOLASAGFYVGRNDYKCCGCGLRWCESGDDPWVEHAKWPPRC 48
|||:||||:| | | | | | | | | | | | | | | | | | | |
DB 105 PELLAAGFFHTGHODKVCFCGGLQSWKRDDPWTEHAKWPPSCQ 152

RESULT 12
Q9H2A8 PRELIMINARY; PRT; 298 AA.
ID Q9H2A8;
AC Q9H2A8;
DT 01-MAR-2001 (TREMblrel. 16, Created)
DT 01-MAR-2001 (TREMblrel. 16, Last sequence update)
DT 01-JUN-2001 (TREMblrel. 17, Last annotation update)
DE INHIBITOR OF APOPTOSIS PROTEIN KIAA (BA261N1.1.2) (BACULOVIRAL IAP
REPEAT-CONTAINING PROTEIN 7 (LIVIN), ISOFORM 2).
GN Birc7.
OS Homo sapiens (Human).
OC Eukaryota; Metazoa; Chordata; Vertebrata; Euteleostomi;
OC Mammalia; Eutheria; Primates; Catarrhini; Homiidae; Homo.
OX NCBI_Taxid=9606;
RN [1]
RP SEQUENCE FROM N.A.
RA TISSUE-KIDNEY;
RC Lin J.-H., Deng G., Huang Q., Morser J.;
RT "A Novel member of the inhibitor of apoptosis protein family.";
RL Biochem. Biophys. Res. Commun. 0:0-0(2000).
[12]
RP SEQUENCE FROM N.A.
RA Ashab Y., Allan A., Pollack A., Panet A., Ben-Yehuda D.;
RT "Two splicing variants of a new inhibitor of apoptosis gene with
RL different biological properties and tissue distribution pattern.";
RN Submitted (Apr-2001) to the EMBL/GenBank/DBJ databases.
[13]
RP SEQUENCE FROM N.A.
RA Hall R.;
RL Submitted (Apr-2001) to the EMBL/GenBank/DBJ databases.
CC -1- SIMILARITY: CONTAINS A RING-TYPE ZINC FINGER.
DR EMBL; AF301009; AAC37878.1; -
DR EMBL; AJ309298; CAC37338.1; -
DR EMBL; AL121827; CAC36111.1; -
DR InterPro: IPR001370; BIR.
DR InterPro: IPR001841; Znf_fing.
DR Pfam: PF00653; BIR; 1.
DR Pfam: PF00097; zf-C3HC4; 1.

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DR SMART; SM00238; BIR; 1.
DR SMART; SM00184; RING; 1.
DR PROSITE; PS50143; BIR_REPEAT_2; 1.
DR PROSITE; PS00518; ZINC_FINGER_C3HC4; UNKNOWN_1.
DR Zinc-finger.
SQ SEQUENCE 298 AA; 32798 MW; B2EAAE531BEC101 CRC64;

Query Match 66.1%; Score 195; DB 4; Length 298;
Best Local Similarity 62.5%; Pred. No. 1.9e-17;
Matches 30; Conservative 7; Mismatches 11; Indels 0; Gaps 0;

QY 1 PEOLASAGFYVGRNDYKCCGCGLRWCESGDDPWVEHAKWPPRC 48
|||:||||:| | | | | | | | | | | | | | | | | | | |
DB 105 PELLAAGFFHTGHODKVCFCGGLQSWKRDDPWTEHAKWPPSCQ 152

RESULT 13
Q9YNL8 PRELIMINARY; PRT; 281 AA.
ID Q9YNL8;
AC Q9YNL8;
DT 01-MAY-1999 (TREMblrel. 10, Created)
DT 01-MAY-1999 (TREMblrel. 10, Last sequence update)
DT 01-JUN-2001 (TREMblrel. 17, Last annotation update)
DE IAP PROTEIN.
GN IAP.
OS Choriostoneura fumiferana nuclear polyhedrosis virus (CFMNPV).
OC Viruses; dsDNA viruses, no RNA stage; Baculoviridae;
OC Nucleopolydrovirus.
OX NCBI_Taxid=10448;
RN [1]
RP SEQUENCE FROM N.A.
RA STRAIN=IRELAND;
RC Lauzon H., Arif B.M., Ladd T., Pall R.;
RT "cfMNPV IAP gene.";
RL Submitted (Dec-1996) to the EMBL/GenBank/DBJ databases.
CC -1- SIMILARITY: CONTAINS A RING-TYPE ZINC FINGER.
DR EMBL; U82510; AAD00537.1; -
DR HSSP; Q13490; IOBH.
DR InterPro: IPR001370; BIR.
DR InterPro: IPR001841; Znf_fing.
DR Pfam: PF00653; BIR; 2.
DR Pfam: PF00097; zf-C3HC4; 1.
DR SMART; SM00238; BIR; 2.
DR SMART; SM00184; RING; 1.
DR PROSITE; PS01282; BIR_REPEAT_1; UNKNOWN_1.
DR PROSITE; PS50143; BIR_REPEAT_2; 2.
DR Zinc-finger.
SQ SEQUENCE 281 AA; 32090 MW; B2D9BE8A359F105E CRC64;

Query Match 65.1%; Score 192; DB 12; Length 281;
Best Local Similarity 66.0%; Pred. No. 4.3e-17;
Matches 31; Conservative 5; Mismatches 11; Indels 0; Gaps 0;

QY 1 PEOLASAGFYVGRNDYKCCGCGLRWCESGDDPWVEHAKWPPRC 47
|||:||||:| | | | | | | | | | | | | | | | | | | |
DB 142 PELLAAGFFHTGRSUKVCFYCDGGLDNWDDDEPMDGHALMWGRC 188

RESULT 14
Q9U492 PRELIMINARY; PRT; 379 AA.
ID Q9U492;
AC Q9U492;
DT 01-MAY-2000 (TREMblrel. 13, Created)
DT 01-MAY-2000 (TREMblrel. 13, Last sequence update)
DT 01-JUN-2001 (TREMblrel. 17, Last annotation update)
DE INHIBITOR OF APOPTOSIS PROTEIN.
GN IAP1.
OS Trichoplusia ni (Cabbage looper).
OC Eukaryota; Metazoa; Arthropoda; Tracheata; Hexapoda; Insecta;
OC Pterygota; Neoptera; Endopterygota; Lepidoptera; Glossata; Ditrysia;
OC Noctuidae; Noctuidae; Plusiinae; Trichoplusia.

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Search completed: January 7, 2002, 16:03:54
 Job time: 1413 sec

OX NCBI_Taxid-7111;
 RN {}
 RP SEQUENCE FROM N.A.
 RX MEDLINE-20052908; PubMed-10593985;
 RA Seshagiri S., Vucic D., Lee J., Dixit V.M.;
 RT "Baculovirus-based genetic screen for antiapoptotic genes identifies a
 novel IAP.";
 RL J. Biol. Chem. 274:36769-36773(1999).
 DR EMBL: AF195528; AAF19819.1; -.
 DR HSSP: Q13490.10BH.
 DR InterPro: IPR001370; BIR.
 DR InterPro: IPR001841; znf_ling.
 DR Pfam: PF00653; BIR; 2.
 DR SMART: SM00238; BIR; 2.
 DR SMART: SM00184; RING; 1.
 DR PROSITE: PS01282; BIR_REPEAT_1; UNKNOWN_2.
 DR PROSITE: PS50143; BIR_REPEAT_2; 2.
 SO SEQUENCE 379 AA; 41857 MW; 069381A012D9DE65 CRC64;

Query Match 64.1%; Score 189; DB 5; Length 379;
 Best Local Similarity 61.7%; Pred. No. 1.5e-16;
 Matches 29; Conservative 7; Mismatches 11; Indels 0; Gaps 0;

OY 1 PEQLASAGFYVGRNDVVKFCFCGCGSLRCWESGDDPWWEHAKWPPRC 47
 ||:|||||:| ||| |||||: ||: ||: ||: |||
 Db 227 PEELAEAGFYTGQDKTKCFYCDGSLKDMENDVPMEOHAKWPPRC 273

RESULT 15
 Q9NJ07 PRELIMINARY; PRT; 377 AA.

AC Q9NJ07;
 DT 01-OCT-2000 (TrEMBLrel. 15, Created)
 DT 01-OCT-2000 (TrEMBLrel. 15, last sequence update)
 DT 01-JUN-2001 (TrEMBLrel. 17, last annotation update)
 DE INHIBITOR OF APOPTOSIS PROTEIN.
 GN IAP.
 OS Spodoptera frugiperda (fall armyworm).
 OC Eukaryota; Metazoa; Arthropoda; Tracheata; Hexapoda; Insecta;
 OC Pterygota; Neoptera; Endopterygota; Lepidoptera; Glossata; Dileptysia;
 OC Noctuoidea; Noctuidae; Amphipyrinae; Spodoptera.
 OX NCBI_Taxid-7108;
 RN {}
 RP SEQUENCE FROM N.A.
 RX MEDLINE-20144055; PubMed-10677478;
 RA Huang Q., Devereux Q.L., Maeda S., Salvesen G.S., Stenmick H.R.,
 RA Hammock B.D., Reed J.C.;
 RT "Evolutionary conservation of apoptosis mechanisms: Lepidopteran and
 RT baculoviral inhibitor of apoptosis proteins are inhibitors of
 RT mammalian caspase-9.";
 RL Proc. Natl. Acad. Sci. U.S.A. 97:1427-1432(2000).
 DR EMBL: AF166378; AAF35285.1; -.
 DR InterPro: IPR001370; BIR.
 DR InterPro: IPR001841; znf_ling.
 DR Pfam: PF00653; BIR; 2.
 DR SMART: SM00238; BIR; 2.
 DR SMART: SM00184; RING; 1.
 DR PROSITE: PS01282; BIR_REPEAT_1; UNKNOWN_2.
 DR PROSITE: PS50143; BIR_REPEAT_2; 2.
 SO SEQUENCE 377 AA; 41871 MW; F1207D2D3ED60B3B CRC64;

Query Match 63.7%; Score 188; DB 5; Length 377;
 Best Local Similarity 61.7%; Pred. No. 2e-16;
 Matches 29; Conservative 7; Mismatches 11; Indels 0; Gaps 0;

OY 1 PEQLASAGFYVGRNDVVKFCFCGCGSLRCWESGDDPWWEHAKWPPRC 47
 ||:|||||:| ||| |||||: ||: ||: ||: |||
 Db 228 PEELAEAGFYTGQDKTKCFYCDGSLKDMENDVPMEOHAKWPPRC 274
